

RAW SEQUENCE LISTING ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 09/539,992ASource: -/600Date Processed by STIC: $-5/7/2\infty2$

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.
PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRF SUBMISSION QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216. PATENTIN 2.1 e-mail help: patin21help@uspto.gov or phone 703-306-4119 (R. Wax) PATENTIN 3.0 e-mail help: patin3help@uspto.gov or phone 703-306-4119 (R. Wax)

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 3.1 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:

http://www.uspto.gov/web/offices/pac/checker

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail.

Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom.

Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

- 1. EFS-Bio (<http://www.uspto.gov/ebc/efs/downloads/documents.htm>, EFS Submission User Manual ePAVE)
- 2. U.S. Postal Service: U.S. Patent and Trademark Office, Box Sequence, P.O. Box 2327, Arlington, VA 22202
- 3. Hand Carry directly to:
 U.S. Patent and Trademark Office, Technology Center 1600, Reception Area, 7th Floor, Examiner Name, Sequence Information, Crystal Mall One, 1911 South Clark Street, Arlington, VA 22202

 Or
 - U.S. Patent and Trademark Office, Box Sequence, Customer Window, Lobby, Room 1B03, Crystal Plaza Two, 2011 South Clark Place, Arlington, VA 22202
- 4. Federal Express, United Parcel Service, or other delivery service to: U.S. Patent and Trademark Office, Box Sequence, Room 1B03-Mailroom, Crystal Plaza Two, 2011 South Clark Place, Arlington, VA 22202

Revised 01/29/2002



MAY 2 3 2002

TECh UENTER 1600/2900



RAW SEQUENCE LISTING DATE: 05/16/2002 PATENT APPLICATION: US/09/539,992A TIME: 18:37:12

```
Does Not Comply
                     Input Set : A:\pto.vsk.txt
                                                                     Corrected Diskette Needed
                     Output Set: N:\CRF3\05162002\I539992A.raw
      3 <110> APPLICANT: Council of Scientific and Industrial Research
      5 <120> TITLE OF INVENTION: A new and distinct somaclonal variety of Rose Scented
Geranium "Parimal".
      7 <130> FILE REFERENCE: 345
      9 <140> CURRENT APPLICATION NUMBER: US 09/539,992A
     10 <141> CURRENT FILING DATE: 2000-03-30
     12 <160> NUMBER OF SEQ ID NOS: 13
     14 <170> SOFTWARE: PatentIn version 3.1
     16 <210> SEQ ID NO: 1
    17 <211> LENGTH: 10
     18 <212> TYPE: DNA
    19 <213> ORGANISM: Artificial sequence
    21 <220> FEATURE:
     22 <223> OTHER INFORMATION: Primers for generating RAPD profile of Plant 'Parimal'.
     24 <400> SEQUENCE: 1
     25 gtgcaatcag
                                                                               10
     28 <210> SEQ ID NO: 2
     29 <211> LENGTH: 10
     30 <212> TYPE: DNA
     31 <213> ORGANISM: artificial sequence
     33 <220> FEATURE:
     34 <223> OTHER INFORMATION: Primers for generating RAPD profile of Plant 'Parimal'.
```

- 36 <400> SEOUENCE: 2
- 37 aggatacgtg
- 40 <210> SEQ ID NO: 3
- 41 <211> LENGTH: 10
- 42 <212> TYPE: DNA
- 43 <213> ORGANISM: Artificial sequence
- 45 <220> FEATURE:
- 46 <223> OTHER INFORMATION: Primers for generating RAPD profile of Plant 'Parimal'.
- 48 <400> SEOUENCE: 3
- 49 aagatagcgg

10

10

- 52 <210> SEQ ID NO: 4
- 53 <211> LENGTH: 10
- 54 <212> TYPE: DNA
- 55 <213> ORGANISM: Artificial Sequence
- 57 <220> FEATURE:
- 58 <223> OTHER INFORMATION: Primers for generating RAPD profile of Plant 'Parimal'.
- 60 <400> SEQUENCE: 4
- 61 ggatctgaac

10

- 64 <210> SEQ ID NO: 5
- 65 <211> LENGTH: 10
- 66 <212> TYPE: DNA
- 67 <213> ORGANISM: Artificial Sequence

DATE: 05/16/2002

TIME: 18:37:12

Input Set : A:\pto.vsk.txt Output Set: N:\CRF3\05162002\I539992A.raw 69 <220> FEATURE: 70 <223> OTHER INFORMATION: Primers for generating RAPD profile of Plant 'Parimal'. 72 <400> SEQUENCE: 5 73 ttgtctcagg 10 76 <210> SEQ ID NO: 6 77 <211> LENGTH: 10 78 <212> TYPE: DNA 79 <213> ORGANISM: Artificial Sequence 81 <220> FEATURE: 82 <223> OTHER INFORMATION: Primers for generating RAPD profile of Plant 'Parimal'. 84 <400> SEQUENCE: 6 85 gtcctactcg 10 88 <210> SEQ ID NO: 7 89 <211> LENGTH: 10 90 <212> TYPE: DNA C--> 91 <213> ORGANISM: Artificial Sequence) W--> 93 $\angle 220$ FEATURE: $W--> 93(\langle 223 \rangle)$ OTHER INFORMATION: 93 <400 SEQUENCE: 7 94 tgcgcgatcg 10 97 <210> SEQ ID NO: 8 98 <211> LENGTH: 10 99 <212> TYPE: DNA 100 <213> ORGANISM: Artificial Sequence 102 <220> FEATURE: 103 <223> OTHER INFORMATION: Primers for generating RAPD profile of Plant 'Parimal'. 105 <400> SEQUENCE: 8 106 aacgtacgcg 10 109 <210> SEQ ID NO: 9 110 <211> LENGTH: 10 111 <212> TYPE: DNA 112 <213> ORGANISM: Artificial Sequence 114 <220> FEATURE: 115 <223> OTHER INFORMATION: Primers for generating RAPD profile of Plant 'Parimal'. 117 <400> SEQUENCE: 9 118 gcacgccgga 10 121 <210> SEQ ID NO: 10 122 <211> LENGTH: 10 123 <212> TYPE: DNA 124 <213 ORGANISM: Artificial Sequence -) see p. 4 W--> 126 <220> FEATURE: ____ W--> $126 (\langle 223 \rangle)$ OTHER INFORMATION: 126 **₹400**> SEQUENCE: 10 127 caccetgcgc 10 130 <210> SEQ ID NO: 11 131 <211> LENGTH: 10 132 <212> TYPE: DNA 133 <212> ORGANISM: Artificial Sequence W--> 135 (220) FEATURE: see p. 4

RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/539,992A

10

RAW SEQUENCE LISTING DATE: 05/16/2002 TIME: 18:37:12 PATENT APPLICATION: US/09/539,992A

Input Set : A:\pto.vsk.txt

Output Set: N:\CRF3\05162002\I539992A.raw

W--> 135 <223> OTHER INFORMATION: 135 <400> SEQUENCE: 11 136 ggactccacg 10 139 <210> SEQ ID NO: 12 140 <211> LENGTH: 10 141 <212> TYPE: DNA 142 <213> ORGANISM: Artificial Sequence 144 <220> FEATURE: 145 <223> OTHER INFORMATION: Primers for generating RAPD profile of Plant 'Parimal'. 147 <400> SEQUENCE: 12 148 aagatagcgg 10 151 <210> SEQ ID NO: 13 152 <211> LENGTH: 10 153 <212> TYPE: DNA 154 <213> ORGANISM: Artificial Sequence 156 <220> FEATURE: 157 <223> OTHER INFORMATION: Primers for generating RAPD profile of Plant 'Parimal'.

159 <400> SEQUENCE: 13

160 ctatcgccgc

RAW SEQUENCE LISTING ERROR SUMMARY PATENT APPLICATION: US/09/539,992A

DATE: 05/16/2002 TIME: 18:37:13

Input Set : A:\pto.vsk.txt

Output Set: N:\CRF3\05162002\I539992A.raw

Invalid Line Length:

The rules require that a line not exceed 72 characters in length. This includes spaces.

Seq#:1; Line(s) 5

Use of <220> Feature(NEW RULES):

Sequence(s)_are missing the <220> Feature and associated headings.
Use of <220> to <223> is MANDATORY if <213> ORGANISM is "Artificial Sequence" or "Unknown". Please explain source of genetic material in <220> to <223> section (See "Federal Register," 6/01/98, Vol. 63, No. 104,pp.29631-32) (Sec.1.823 of new Rules)

Seq#:7,10,11

file://C:\Crf3\Outhold\VsrI539992A.htm

VERIFICATION SUMMARY

DATE: 05/16/2002

PATENT APPLICATION: US/09/539,992A

TIME: 18:37:13

Input Set : A:\pto.vsk.txt

Output Set: N:\CRF3\05162002\I539992A.raw

L:91 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:7

L:93 M:258 W: Mandatory Feature missing, <220> FEATURE:

L:93 M:258 W: Mandatory Feature missing, <223> OTHER INFORMATION:

L:126 M:258 W: Mandatory Feature missing, <220> FEATURE:

L:126 M:258 W: Mandatory Feature missing, <223> OTHER INFORMATION:

L:135 M:258 W: Mandatory Feature missing, <220> FEATURE:

L:135 M:258 W: Mandatory Feature missing, <223> OTHER INFORMATION: